

SEQUENCE LISTING

<110> BAKALETZ, et al.

<120> NONTYPEABLE HAEMOPHILUS INFLUENZAE VIRULENCE FACTORS

<130> 28335/39196A

<140> To be assigned

<141> Herewith

<150> US 60/458,234

<151> 2003-03-27

<160> 41

<170> PatentIn version 3.2

<210> 1

<211> 1695

<212> DNA

<213> H. influenzae

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cgtatttttc atcaacgagc agaatatatt ttgaatcaaa atttttctaa agtagcgaca	660
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<212> DNA

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ctttccagta tgaaccaaaa tcagggaaaca acaattttac ttacgagtaa cgatattaaa 660
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ccacgtttaa atattggaca aatatttagac gaaccattaa gcctagcgac agattggaca 360
gaaacacaac gtaatgaaaa aatcttgag accctctctc ttgttgact ttatcctgat 420
tacacaaatc tcaatattaa gcatctctc atcagccaaa agcagcgggt tgccctagca 480
cgcgcattaa ttttagcacc agaaattatt ataatagatg atgcaattgg caatttagat 540
gcttctgtac gtattcaatt gcttaattt acccttgatt tacaacaacg ttttaggtata 600
tcttatattt atgtggaca ggtctcggt gtaattaaac atattgcaga tacgattatc 660
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gcttggtaa aagacaaaaa cactcactaa 810

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 <213> Homo sapiens

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30	

Glu Asn Gly Leu Thr Tyr Cys Thr His Ala Ser Gly Phe Ser Phe Asn	
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45	

Pro Gln Thr Ala Asp Ala Gly Thr Ser Met Asn Val Val Thr Glu Gln	
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60	

Ile Tyr Asn Lys Leu Phe Asp Ile Lys Asn His Ser Ala Thr Leu Thr	
65	70
75	80

Pro Met Leu Ala Gln Ser Tyr Ser Ile Ser Ala Asp Gly Lys Glu Ile	
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95	

Leu Leu Asn Leu Arg His Gly Val Lys Phe His Gln Thr Pro Trp Phe	
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110	

Thr Pro Thr Arg Asp Phe Asn Ala Glu Asp Val Val Phe Ser Ile Asn	
115	120
125	

Arg Val Leu Gly His Asn Thr Tyr Leu Pro Thr Leu Ala Glu Ala Asn
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Val Thr Tyr Ser Asn Pro Gln Tyr Arg Val Phe His Glu Gln Ala Arg
145 150 155 160

Lys Val Arg Phe Pro Tyr Phe Asp Ser Ile Lys Leu Asn Glu Lys Ile
165 170 175

Lys Ser Val Thr Ala Leu Ser Pro Tyr Gln Val Lys Ile Glu Leu Phe
180 185 190

Ala Pro Asp Ser Ser Ile Leu Ser His Leu Ala Ser Gln Tyr Ala Ile
195 200 205

Ile Phe Ser Gln Glu Tyr Ala Tyr Gln Leu Ser Ala Asp Asp Asn Leu
210 215 220

Ala Gln Leu Asp Thr His Pro Val Gly Thr Gly Pro Tyr Gln Val Lys
225 230 235 240

Asp Tyr Val Tyr Asn Gln Tyr Val Arg Leu Val Arg Asn Glu Asn Tyr
245 250 255

Trp Lys Lys Glu Ala Lys Ile Glu His Ile Ile Val Asp Leu Ser Thr
260 265 270

Asp Arg Ser Gly Arg Leu Val Lys Phe Phe Asn Asn Glu Cys Gln Ile
275 280 285

Ala Ser Tyr Pro Glu Val Ser Gln Ile Gly Leu Leu Lys Asn Asp Asp
290 295 300

Lys His Tyr Tyr Met Gln Ser Thr Asp Gly Met Asn Leu Ala Tyr Leu
305 310 315 320

Ala Phe Asn Phe Asp Lys Pro Leu Met Arg Asp His Glu Ile Arg Ala
325 330 335

Ala Ile Ser Gln Ser Leu Asn Arg Ala Arg Ile Ile His Ser Ile Tyr
340 345 350

His Asn Thr Ala Thr Val Ala Asn Asn Ile Ile Pro Glu Val Ser Trp
355 360 365

Ala Ser Thr Val Asn Thr Pro Glu Phe Glu Phe Asp Tyr His Pro Lys
370 375 380

Ile Ala Lys Asn Lys Leu Ala Asp Lys Asn Leu Leu Leu Asn Leu Trp
385 390 395 400

Val Ile Asn Glu Glu Gln Val Tyr Asn Pro Ala Pro Phe Lys Met Ala
405 410 415

Glu Met Ile Lys Trp Asp Leu Ala Gln Ala Gly Val Lys Val Lys Val
420 425 430

Arg Ala Val Thr Arg Pro Phe Leu Thr Ala Gln Leu Arg Asn Gln Ser
435 440 445

Glu Asn Tyr Asp Leu Ile Leu Ser Gly Trp Leu Ala Gly Asn Leu Asp
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Pro Asp Gly Phe Met Arg Pro Ile Leu Ser Cys Gly Thr Lys Asn Glu
465 470 475 480

Leu Thr Asn Leu Ser Asn Trp Cys Asn Glu Glu Phe Asp Gln Phe Met
485 490 495

Asp Arg Ala Ile Thr Thr Ser His Leu Ser Ser Arg Ala Lys Ala Tyr
500 505 510

Asn Glu Ala Gln Glu Leu Val Leu Arg Glu Leu Pro Ile Ile Pro Ile
515 520 525

Ala Asn Val Lys Arg Ile Leu Val Ala Asn Ser Arg Val Lys Gly Val
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Lys Met Thr Pro Phe Gly Ser Leu Asp Phe Ser Thr Leu Tyr Phe Ile
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Gln Glu Lys His

<210> 8
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<212> PRT
<213> H. influenzae

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Asn	Ala	Asn	Leu	Val	Thr	Gln	Asn	Ile	Tyr	Ile	Gly	Tyr	Phe	His	Tyr
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Leu	Gly	Thr	Leu	Leu	Gln	Gly	Asp	Phe	Gly	Ile	Thr	Tyr	Asn	Gly	Gly
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Lys	Ser	Leu	Met	Asn	Leu	Ile	Leu	Thr	Val	Leu	Pro	Pro	Thr	Leu	Glu
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Leu	Cys	Phe	Ile	Thr	Leu	Phe	Leu	Ala	Phe	Ile	Phe	Gly	Leu	Pro	Leu
														85	90
Gly	Ile	Ile	Ser	Ala	Val	Asn	Ser	Glu	Gln	Val	Phe	Ala	Lys	Ser	Leu
														100	105
Gln	Ile	Leu	Ser	Tyr	Val	Gly	Leu	Ser	Ile	Pro	Ile	Phe	Trp	Leu	Ala
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Pro	Ile	Leu	Leu	Tyr	Val	Ala	Ala	Leu	Ser	His	Trp	Glu	Ile	Ala	Ala
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Ile	Gly	Gln	Tyr	Asn	Leu	Leu	Tyr	Glu	Ile	Lys	Pro	Ile	Thr	Gly	Phe
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Pro	Val	Ile	Asp	Met	Trp	Phe	Met	Glu	Val	Pro	Tyr	Arg	Thr	Lys	Ile
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Val	Gln	Asn	Ile	Leu	Gln	His	Leu	Ala	Leu	Pro	Thr	Leu	Val	Leu	Cys
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Ile	Leu	Pro	Thr	Met	Glu	Ile	Ile	Arg	Ile	Ile	His	Gln	Arg	Ala	Glu
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Tyr	Ile	Leu	Asn	Gln	Asn	Phe	Ser	Lys	Val	Ala	Thr	Thr	Arg	Gly	Trp
														210	215
Ser	Lys	Trp	Lys	Ile	Leu	His	Gln	Tyr	Val	Phe	Arg	Asn	Thr	Phe	Pro
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Leu	Leu	Val	Pro	Gln	Val	Pro	Arg	Val	Phe	Thr	Leu	Val	Leu	Thr	Gln
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Cys	Met	Leu	Val	Glu	Thr	Ala	Leu	Gly	Trp	Pro	Gly	Ile	Gly	Arg	Trp
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Leu	Ile	Asn	Ala	Val	Asn	Glu	Gln	Asp	Tyr	Asn	Ser	Ile	Ala	Ala	Gly
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														285	

Val	Ile	Val	Ile	Gly	Val	Cys	Ile	Ile	Leu	Ile	Asp	Thr	Phe	Thr	Lys
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20			25								30				
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35					40							45			
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Thr	Leu	Gly	Ser	Ala	Leu	Leu	Val	Val	Phe	Ser	Val	Ala	Ile	Ile	Gly
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Gly	Ala	Leu	Gly	Ile	Ile	Ala	Gly	Leu	Leu	Lys	Gly	Ile	Lys	Ala	Arg
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165						170						175			
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Lys Leu Glu Gly Ile Ser Asn Gln Thr Leu Leu Lys Ser Thr Ile Leu
195 200 205

Pro Asn Ile Thr Val Ile Tyr Ile Gln Glu Val Ala His Ala Phe Val
210 215 220

Ile Ala Val Leu Asp Ile Ser Ala Leu Ser Phe Ile Ser Leu Gly Ala
225 230 235 240

Gln Arg Pro Thr Pro Glu Trp Gly Ala Met Ile Lys Asp Ser Leu Glu
245 250 255

Leu Leu Tyr Leu Ala Pro Trp Thr Val Leu Leu Pro Gly Phe Ala Ile
260 265 270

Ile Phe Thr Ile Leu Leu Ser Ile Ile Phe Ser Asn Gly Leu Thr Lys
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Ala Ile Asn Gln His Gln Glu
290 295

<210> 10
<211> 349
<212> PRT
<213> H. influenzae

<400> 10

Met Ala Leu Leu Asp Ile Cys Asn Leu Asn Ile Glu Ile Gln Thr Ser
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Asn Gly Arg Ile Lys Ile Val Asp Gly Val Asn Leu Ser Leu Asn Glu
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Gly Glu Ile Ser Gly Leu Val Gly Glu Ser Gly Ser Gly Lys Ser Leu
35 40 45

Ile Ala Lys Val Ile Cys Asn Ala Ile Lys Glu Asn Trp Ile Ile Thr
50 55 60

Ala Asp Arg Phe Arg Phe His Asp Ile Glu Leu Leu Lys Leu Ser Pro
65 70 75 80

Asn Lys Arg Arg Lys Ile Val Gly Lys Glu Ile Ser Met Ile Phe Gln
85 90 95

Asn Pro Leu Ser Cys Leu Asp Pro Ser Arg Lys Ile Gly Lys Gln Leu
100 105 110

Ile Gln Asn Ile Pro Asn Trp Thr Phe Lys Asn Lys Trp Trp Lys Trp
115 120 125

Phe Gly Trp Lys Lys Arg Arg Ala Ile Glu Leu Leu His Arg Val Gly
130 135 140

Ile Lys Asp His Arg Asp Ile Met Ala Ser Tyr Pro Asn Glu Leu Thr
145 150 155 160

Glu Gly Glu Gly Gln Lys Val Met Ile Ala Met Ala Val Ala Asn Gln
165 170 175

Pro Arg Leu Leu Ile Ala Asp Glu Pro Thr Asn Thr Leu Glu Ser Thr
180 185 190

Thr Ala Leu Gln Val Phe Arg Leu Leu Ser Ser Met Asn Gln Asn Gln
195 200 205

Gly Thr Thr Ile Leu Leu Thr Ser Asn Asp Ile Lys Ser Ile Ser Glu
210 215 220

Trp Cys Asp Gln Ile Ser Val Leu Tyr Cys Gly Gln Asn Thr Glu Ser
225 230 235 240

Ala Pro Thr Glu Ile Leu Ile Glu Ser Pro His His Pro Tyr Thr Gln
245 250 255

Ala Leu Ile Asn Ala Val Pro Asp Phe Thr Gln Pro Leu Gly Phe Lys
260 265 270

Thr Lys Leu Gly Thr Leu Glu Gly Thr Ala Pro Ile Leu Glu Gln Met
275 280 285

Pro Ile Gly Cys Arg Leu Gly Pro Arg Cys Pro Phe Ala Gln Lys Lys
290 295 300

Cys Met Glu Lys Pro Arg Arg Leu Lys Ile Lys Gln His Glu Phe Ser
305 310 315 320

Cys His Tyr Pro Ile Asn Leu Arg Glu Lys Asn Phe Lys Glu Lys Thr
325 330 335

Thr Ala Thr Pro Phe Ile Leu Asn Cys Lys Gly Asn Glu
340 345

<210> 11
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<212> PRT

<213> Homo sapiens

<400> 11

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35 40 45

Ser Gly Lys Ser Thr Leu Val Lys Met Ile Ala Gly Ile Ile Pro Pro
50 55 60

Thr Ser Gly Arg Ile Leu Phe Asn Asp Arg Glu Leu Gln Tyr Gln Asp
65 70 75 80

Ala Gln Ser Arg Ala Lys His Ile Arg Met Val Phe Gln Asp Ala Asn
85 90 95

Ser Ala Phe Asn Pro Arg Leu Asn Ile Gly Gln Ile Leu Asp Glu Pro
100 105 110

Leu Ser Leu Ala Thr Asp Trp Thr Glu Thr Gln Arg Asn Glu Lys Ile
115 120 125

Phe Glu Thr Leu Ser Leu Val Gly Leu Tyr Pro Asp Tyr Thr Asn Leu
130 135 140

Asn Ile Lys His Leu Ser Ile Ser Gln Lys Gln Arg Val Ala Leu Ala
145 150 155 160

Arg Ala Leu Ile Leu Ala Pro Glu Ile Ile Ile Asp Asp Ala Ile
165 170 175

Gly Asn Leu Asp Ala Ser Val Arg Ile Gln Leu Leu Asn Leu Thr Leu
180 185 190

Asp Leu Gln Gln Arg Leu Gly Ile Ser Tyr Ile Tyr Val Gly Gln Asp
195 200 205

Leu Gly Val Ile Lys His Ile Ala Asp Thr Ile Ile Val Met Asp Asp
210 215 220

Gly Lys Met Ile Glu Tyr Gly Ser Pro Gln Asn Leu Phe Thr Asp Pro
225 230 235 240

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aatatccatt	atcaacccaaa	atataaacac	ggcggtacac	cgaaattacc	tgatgatatt	1980

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cattcttaca	gtttgtttc	aacgattgat	ccaacgagtt	ttcttcgttt	acaactaaaa	2160
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gatttcacta	tttgccaaa	tactaatcta	aaaccagaga	tagcaaaaac	aaaagaaatt	2280
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 <212> PRT
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<400> 41

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Val	Met	Leu	Gly	Leu	Thr	Ala	Gly	Val	Ala	Tyr	Ala	Ala	Gln	Pro	Thr
20					25								30		

Asn	Gln	Pro	Thr	Asn	Gln	Pro	Thr	Asn	Gln	Pro	Thr	Asn	Gln	Asn	Gly
35				40					45						

Asn	Val	Ser	Glu	Gln	Leu	Glu	Gln	Ile	Asn	Val	Ser	Gly	Ser	Thr	Glu
50				55				60							

Asn	Ser	Asp	Thr	Lys	Thr	Pro	Pro	Lys	Ile	Ala	Glu	Thr	Val	Lys	Thr
65				70				75				80			

Ala	Lys	Thr	Leu	Glu	Arg	Glu	Gln	Ala	Asn	Asn	Ile	Lys	Asp	Ile	Val
85					90							95			

Lys Tyr Glu Thr Gly Val Thr Val Val Glu Ala Gly Arg Phe Gly Gln
100 105 110

Ser Gly Phe Ala Ile Arg Gly Val Asp Glu Asn Arg Val Ala Ile Asn
115 120 125

Ile Asp Gly Leu Arg Gln Ala Glu Thr Leu Ser Ser Gln Gly Phe Lys
130 135 140

Glu Leu Phe Glu Gly Tyr Gly Asn Phe Asn Asn Thr Arg Asn Gly Ala
145 150 155 160

Glu Ile Glu Thr Leu Lys Glu Val Asn Ile Thr Lys Gly Ala Asn Ser
165 170 175

Ile Lys Ser Gly Ser Gly Ser Leu Gly Gly Ser Val Ile Tyr Lys Thr
180 185 190

Lys Asp Ala Arg Asp Tyr Leu Leu Asn Lys Asp Tyr Tyr Val Ser Tyr
195 200 205

Lys Lys Gly Tyr Ala Thr Glu Asn Asn Gln Ser Phe Asn Thr Leu Thr
210 215 220

Leu Ala Gly Arg Tyr Lys Lys Phe Asp Ala Leu Val Val Thr Thr Ser
225 230 235 240

Arg Asn Gly His Glu Leu Glu Asn Tyr Asp Tyr Lys Asn Ala Asn Ser
245 250 255

Leu Thr Gln Gly Lys Lys Arg Glu Lys Ala Asp Pro Tyr Lys Ile Glu
260 265 270

Gln Asp Ser Thr Leu Leu Lys Leu Ser Phe Asn Pro Thr Glu Asn His
275 280 285

Arg Phe Thr Leu Ala Ala Asp Leu Tyr Glu His Arg Ser Arg Gly Gln
290 295 300

Asp Leu Ser Tyr Thr Leu Lys Tyr Gln Lys Thr Asp Pro Asn Leu Leu
305 310 315 320

Glu Val Asp Ser Arg His Thr Asn Asp Lys Thr Lys Arg Arg Asn Ile
325 330 335

Ser Phe Ser Tyr Glu Asn Phe Ser Gln Thr Pro Phe Trp Asp Thr Leu
340 345 350

Lys Ile Thr Tyr Ser Asp Gln Arg Ile Lys Thr Arg Ala Arg Thr Asp
355 360 365

Asp Tyr Cys Asp Ala Gly Val Arg Tyr Cys Glu Gly Thr Ala Asn Pro
370 375 380

Ala Gly Leu Lys Leu Thr Asp Gly Lys Ile Thr Arg Arg Asp Gly Ser
385 390 395 400

Glu Leu Gln Phe Glu Lys Lys Asp Lys Asn Ile Asp Asn Asn Ile Tyr
405 410 415

Asp Phe Asp Lys Phe Ile Asp Thr Asp Asp Arg Val Ile Glu Gly Lys
420 425 430

Leu Gly Leu Arg Arg Ser Ser Gly Thr Trp Tyr Asp Cys Ser Ile Phe
435 440 445

Asp Cys Lys Asp Lys Thr Lys Met Lys Ile Phe Glu Thr Glu His Pro
450 455 460

Tyr Gly Tyr Gly Thr Thr Gly Thr Trp Lys Lys Asp Phe Glu Leu Glu
465 470 475 480

Ile Lys Lys Leu Asn Asp Lys Asn Phe Ala Arg Val Lys Asp Ala Asn
485 490 495

Asn Lys Thr Tyr Ser Ile Leu Pro Ser Ser Pro Gly Tyr Leu Glu Arg
500 505 510

Leu Trp Gln Glu Arg Asp Leu Asp Thr Asn Thr Gln Gln Leu Asn Leu
515 520 525

Asp Leu Thr Lys Asp Phe Lys Thr Trp Arg Val Glu His Asn Leu Gln
530 535 540

Tyr Gly Ser Ser Tyr Asn Thr Thr Met Lys Arg Met Val Asn Arg Ala
545 550 555 560

Gly Asn Asp Ala Ser Asp Val Gln Trp Trp Ala Glu Pro Thr Leu Gly
565 570 575

Tyr Ser Leu Leu Tyr Asp Lys Pro His Thr Cys Lys Thr Ala Tyr Gly
580 585 590

Gly Trp Lys Ala Asn Leu Cys Pro Arg Val Asp Pro Lys Phe Ser Phe
595 600 605

Leu Leu Pro Ile Lys Thr Lys Glu Lys Ser Val Tyr Leu Phe Asp Asn
610 615 620

Val Val Ile Thr Asp Tyr Leu Ser Phe Asp Leu Gly Tyr Arg Tyr Asp
625 630 635 640

Asn Ile His Tyr Gln Pro Lys Tyr Lys His Gly Val Thr Pro Lys Leu
645 650 655

Pro Asp Asp Ile Val Lys Gly Leu Phe Ile Pro Leu Pro Ser Gly Lys
660 665 670

Asn Asn Asn Asp Asp Pro Glu Val Lys Lys Asn Val Gln Gln Asn Ile
675 680 685

Asp Tyr Ile Ala Lys Gln Asn Lys Lys Tyr Lys Ala His Ser Tyr Ser
690 695 700

Phe Val Ser Thr Ile Asp Pro Thr Ser Phe Leu Arg Leu Gln Leu Lys
705 710 715 720

Tyr Ser Lys Gly Phe Arg Ala Pro Thr Ser Asp Glu Met Tyr Phe Thr
725 730 735

Phe Lys His Pro Asp Phe Thr Ile Leu Pro Asn Thr Asn Leu Lys Pro
740 745 750

Glu Ile Ala Lys Thr Lys Glu Ile Ala Phe Thr Leu His Asn Asp Asp
755 760 765

Trp Gly Phe Ile Ser Thr Ser Leu Phe Lys Thr Asn Tyr Lys Asn Phe
770 775 780

Ile Asp Leu Ile Phe Lys Gly Glu Lys Asp Phe Lys Leu Val Ser Gly
785 790 795 800

Gly Ser Thr Leu Pro Phe Ser Leu Tyr Gln Asn Ile Asn Arg Asp Ser
805 810 815

Ala Val Val Lys Gly Ile Glu Ile Asn Ser Lys Val Phe Leu Gly Lys
820 825 830

Met Ala Lys Phe Met Asp Gly Phe Asn Leu Ser Tyr Lys Tyr Thr Tyr
835 840 845

Gln Lys Gly Arg Met Asp Gly Asn Ile Pro Met Asn Ala Ile Gln Pro
850 855 860

Lys Thr Met Val Tyr Gly Leu Gly Tyr Asp His Pro Ser Gln Lys Phe
865 870 875 880

Gly Phe Asn Phe Tyr Thr Thr His Val Ala Ser Lys Asn Pro Glu Asp
885 890 895

Thr Tyr Asp Ile Tyr Ala Lys Asp Lys Asn Gln Thr Asn Thr Ser Ile
900 905 910

Lys Trp Arg Ser Lys Ser Tyr Thr Ile Leu Asp Leu Ile Gly Tyr Val
915 920 925

Gln Pro Ile Lys Asn Leu Thr Ile Arg Ala Gly Val Tyr Asn Leu Thr
930 935 940

Asn Arg Lys Tyr Ile Thr Trp Asp Ser Ala Arg Ser Ile Arg Ser Phe
945 950 955 960

Gly Thr Ser Asn Val Ile Asp Gln Lys Thr Gly Gln Gly Ile Asn Arg
965 970 975

Phe Tyr Ala Pro Gly Arg Asn Tyr Lys Met Ser Val Gln Phe Glu Phe
980 985 990